

## Sequence listing

<110> Epigenomics AG

<120> Method for the analysis of methylation patterns within nucleic acids  
by means of mass spectrometry.

<160> 3

<210> 1

<211> 17

<212> DNA

<213> Homo Sapiens

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<213> Homo Sapiens

<400> 2

- gcttggggga ataggag 18

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<211> 22

<212> DNA

<213> Homo Sapiens

<400> 3

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